

SEQUENCE LISTING

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YATOMI, Takehiro
SUDA, Takashi

<120> PROPHYLACTIC/THERAPEUTIC AGENT

<130> 1110-235P

<140> 09/297,328

<141> 1999-04-29

<150> PCT/JP97/03978

<151> 1997-10-31

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<170> PatentIn Ver. 2.0

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Gly	Thr	Arg	Cys	Asp	Ile	Gln	Met	Thr	Gln	Thr	Thr	Ser	Ser	Leu	Ser	
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gcc	tct	ctg	gga	gac	aga	gtc	acc	atc	agt	tgc	agg	gcc	agt	cag	gac	144
Ala	Ser	Leu	Gly	Asp	Arg	Val	Thr	Ile	Ser	Cys	Arg	Ala	Ser	Gln	Asp	
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Ile	Ser	Asn	Tyr	Leu	Asn	Trp	Tyr	Gln	Gln	Lys	Pro	Asp	Gly	Thr	Val	
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aaa	ctc	ctg	atc	tac	tac	aca	tca	aga	tta	cac	tca	gga	gtc	cca	tca	240
Lys	Leu	Leu	Ile	Tyr	Tyr	Thr	Ser	Arg	Leu	His	Ser	Gly	Val	Pro	Ser	
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Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asn	Tyr	Ser	Leu	Thr	Ile	Ser	
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aac	ctg	gaa	caa	gga	gat	att	gcc	act	tac	ttt	tgc	caa	cag	ggc	agt	336
Asn	Leu	Glu	Gln	Gly	Asp	Ile	Ala	Thr	Tyr	Phe	Cys	Gln	Gln	Gly	Ser	
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 Lys Leu Leu Ile Tyr Tyr Thr Ser Arg Leu His Ser Gly Val Pro Ser
 65 70 75 80
 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asn Tyr Ser Leu Thr Ile Ser
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 Ala Gln Ala Gln Ile Gln Leu Val Gln Ser Gly Pro Glu Leu Lys Lys
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 Pro Gly Glu Thr Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe

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Lys Trp Met Gly Met Ile Tyr Thr Asp Thr Gly Glu Pro Ser Tyr Ala			
65	70	75 80	
gaa gag ttc aag ggg cgg ttt gcc ttc tct ttg gag acc tct gcc agc			288
Glu Glu Phe Lys Gly Arg Phe Ala Phe Ser Leu Glu Thr Ser Ala Ser			
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act gcc tat ttg cag atc aac ttc ctc aaa aat gag gac acg gct aca			336
Thr Ala Tyr Leu Gln Ile Asn Phe Leu Lys Asn Glu Asp Thr Ala Thr			
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Tyr Phe Cys Val Arg Phe Tyr Trp Asp Tyr Phe Asp Tyr Trp Gly Gln			
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Thr Ala Tyr Leu Gln Ile Asn Phe Leu Lys Asn Glu Asp Thr Ala Thr			
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Gly Ser Thr Gly Asp Ile Gln Met Thr Gln Ser Pro Ser Thr Leu Ser
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gct agc gtc ggg gat agg gtc acc ata act tgc agg gca agt cag gac 144
Ala Ser Val Gly Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Asp
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Ile Ser Asn Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro
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Lys Leu Leu Ile Tyr Tyr Thr Ser Arg Leu His Ser Gly Val Pro Ser
65 70 75 80
cgc ttc agt ggc agt gga tct ggg acc aat tat acc ctc aca atc tcg 288
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asn Tyr Thr Leu Thr Ile Ser
85 90 95
agt ctg cag cca gat gat ttc gcc act tat ttt tgc caa cag ggt agt 336
Ser Leu Gln Pro Asp Asp Phe Ala Thr Tyr Phe Cys Gln Gln Gly Ser
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Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	Leu	Met	Ile				
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gac	cct	gag	gtc	aag	ttc	aac	tgg	tac	gtg	gac	ggc	gtg	gag	gtg	cat	677			
Asp	Pro	Glu	Val	Lys	Phe	Asn	Trp	Tyr	Val	Asp	Gly	Val	Glu	Val	His				
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aat	gcc	aag	aca	aag	ccg	cgg	gag	gag	cag	tac	aac	agc	acg	tac	cgt	725			
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Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys	
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gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc ccc atc gag	821
Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu	
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aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gaa cca cag gtg tac	869
Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr	
265 270 275	
acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag gtc agc ctg	917
Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu	
280 285 290	
acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc gtg gag tgg	965
Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp	
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gag agc aat ggg cag ccg gag aac aac tac aag acc acg cct ccc gtg	1013
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Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp	
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Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His	
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Leu Leu Pro Leu Val Leu Thr Ser Val Ala Thr Gln Asn Leu Glu Gly
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Leu His His Asp Gly Gln Phe Cys His Lys Pro Cys Pro Pro Gly Glu
25 30 35

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Arg Lys Ala Arg Asp Cys Thr Val Asn Gly Asp Glu Pro Asp Cys Val
40 45 50

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Pro Cys Gln Glu Gly Lys Glu Tyr Thr Asp Lys Ala His Phe Ser Ser
55 60 65 70

aaa tgc aga aga tgt aga ttg tgt gat gaa gga cat ggc tta gaa gtg 293
Lys Cys Arg Arg Cys Arg Leu Cys Asp Glu Gly His Gly Leu Glu Val
75 80 85

gaa ata aac tgc acc cgg acc cag aat acc aag tgc aga tgt aaa cca 341
Glu Ile Asn Cys Thr Arg Thr Gln Asn Thr Lys Cys Arg Cys Lys Pro
90 95 100

aac ttt ttt tgt aac tct act gta tgt gaa cac tgt gac cct tgc acc 389
Asn Phe Phe Cys Asn Ser Thr Val Cys Glu His Cys Asp Pro Cys Thr
105 110 115

aaa tgt gaa cat gga atc atc aag gaa tgc aca ctc acc agc aac acc 437
Lys Cys Glu His Gly Ile Ile Lys Glu Cys Thr Leu Thr Ser Asn Thr
120 125 130

aag tgc aaa gag gaa gga tcc aga tct aac gag ccc aaa tct tgt gac 485
Lys Cys Lys Glu Glu Gly Ser Arg Ser Asn Glu Pro Lys Ser Cys Asp
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 Asp Glu Pro Asp Cys Val Pro Cys Gln Glu Gly Lys Glu Tyr Thr Asp
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 Lys Ala His Phe Ser Ser Lys Cys Arg Arg Cys Arg Leu Cys Asp Glu
 65 70 75 80
 Gly His Gly Leu Glu Val Glu Ile Asn Cys Thr Arg Thr Gln Asn Thr
 85 90 95
 Lys Cys Arg Cys Lys Pro Asn Phe Phe Cys Asn Ser Thr Val Cys Glu
 100 105 110
 His Cys Asp Pro Cys Thr Lys Cys Glu His Gly Ile Ile Lys Glu Cys
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 145 150 155

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